

Nuclear Norm Subspace Identification (N2SID) for short data batches^{*}

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Abstract: Subspace identification is revisited in the scope of nuclear norm minimization methods. It is shown that essential structural knowledge about the unknown data matrices in the *data equation* that relates Hankel matrices constructed from input and output data can be used in the first step of the numerical solution presented. The structural knowledge comprises the low rank property of a matrix that is the product of the extended observability matrix and the state sequence and the Toeplitz structure of the matrix of Markov parameters (of the system in innovation form). The new subspace identification method is referred to as the N2SID (*twice* the N of Nuclear Norm and SID for Subspace IDentification) method. In addition to include key structural knowledge in the solution it integrates the subspace calculation with minimization of a classical prediction error cost function. The nuclear norm relaxation enables us to perform such integration while preserving *convexity*. The advantages of N2SID are demonstrated in a numerical *open- and closed-loop* simulation study. Here a comparison is made with another widely used SID method, i.e. N4SID. The comparison focusses on the identification with short data batches, i.e. where the number of measurements is a small multiple of the system order.

Keywords: Subspace system identification, Nuclear norm optimization, Rank constraint, Short data batches

1. INTRODUCTION

System identification is a key problem in a large number of scientific areas. Generally there are two families of system identification methods: (1) prediction error methods and (2) subspace methods, Ljung [1999], Verhaegen and Verdult [2007]. Either of these approaches can be treated in the time or frequency domain, and for the sake of simplicity we restrict ourselves to the time domain in this paper.

The central theme in prediction error methods is to parametrize the predictor (observer) to generate an estimate of the output and then formulate an optimization problem to minimize a (weighted) cost function defined on the difference between the measured output and the observer predicted output. This cost function generally is a sample average (for the finite data length case) of the trace of the covariance matrix of the prediction error. Though the prediction error framework provides a vast amount of insights in studying and analyzing the estimated predictor, its main drawback is the non-convexity for general multi-variable state space models in innovation form, as consid-

ered in this paper. The lack of convexity can result in that the optimization method get stuck in a local minimum, and thereby complicating the analysis of the numerical results, such as e.g. the difficulty to distinguish between a bad model estimate due to a local minimum or due to a bad model parametrization. This parametrization needs to be given before starting the parameter optimization problem, and thus the use of the approach can be quite complex and labor intensive for the non-expert user. However, the latter fact has been greatly relieved by computer added software packages such as in Ljung [2007].

Motivated by the drawbacks of prediction error methods, the goal with subspace identification methods is to derive *approximate* models rather than models that are “optimal” with respect to a chosen cost function. The approximation is based on linear algebra transformations and factorizations with structured Hankel matrices constructed from the input-output data. All existing subspace identification methods aim to derive a low rank matrix from which key subspaces, hence the name subspace identification, are derived. The low rank approximation is in general done using a *singular value decomposition* (SVD). Recently a new family of subspace identification methods was presented that use the *nuclear norm* instead of a SVD in order to improve the low rank approximation, Liu and Vandenberghe [2009a,b], Mohan and Fazel [2010], Fazel et al. [2012], Hansson et al. [2012], Liu et al. [2013].

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The drawback of Subspace Identification (SID) is two-fold. First general subspace identification methods lack an optimization criterion in the calculation of the predictor. This drawback is relaxed in a number of recent nuclear norm based SID methods that regularize the low rank approximation problem with a sample average of the trace of the covariance matrix of the prediction error. The second drawback is that the low rank approximation is not performed in the first step of the algorithm. As a consequence, this low rank decomposition does not operate with the original input-output data but with approximate processed data. The latter approximation destroys the low rank property and especially for short data batches, since most of the SID schemes only provide consistent estimates.

As it is known that exploiting structure (that is present in the model) is beneficial, especially when working with short data batches, we review the derivation of subspace identification in order to be able to deal with two key structure properties in the data equation used by SID methods. The data equation is a relationship between the Hankel matrices constructed from the input and output data, respectively. The key structural properties are the low rank property and the Toeplitz structure of unknown model dependent matrices in the data equation. The key in the derivation of the new scheme is that both these structural properties are invoked in the first step of the algorithm. The algorithm is abbreviated by N2SID, standing for Nuclear Norm (two times N, i.e. N2) Subspace Identification.

The paper is organized as follows. In Section 2 the identification problem for identifying a multivariable state space model in a subspace context while taking a prediction error cost function into consideration is presented. The problem formulation does however not require a parametrization of the system matrices of the state space model as is classically done in prediction error methods for parametric input-output models, Ljung [1999]. The key data equation in the analysis and description of subspace identification methods is presented in Section 3. Here we also highlight the important structural properties in the submatrices of this equation, the low rank property of the matrix that is the product of the extended observability matrix and the state sequence of the underlying state space model (given in innovation form) and the (lower triangular) Toeplitz structure of the model Markov parameters. A convex relaxation is presented in Section 4 to take these structural constraints into consideration. Some preliminary results of the performances are illustrated in Section 5 in a comparison study of N2SID with N4SID for both open and closed-loop identification problems with short data batches. Here we consider the identification of second order systems with just 50 data points. Finally, we end this paper with some concluding remarks.

2. THE SUBSPACE IDENTIFICATION PROBLEM

In system identification a challenging problem is to identify Linear Time Invariant (LTI) systems with multiple inputs and multiple outputs using short length data sequences. Taking process and measurement noise into consideration, the model of the LTI system can be written in the following innovation form:

$$\begin{cases} x(k+1) = Ax(k) + Bu(k) + Ke(k) \\ y(k) = Cx(k) + Du(k) + e(k) \end{cases} \quad (1)$$

with $x(k) \in \mathbb{R}^n, y(k) \in \mathbb{R}^p, u(k) \in \mathbb{R}^m$ and $e(k)$ a zero-mean white noise sequences with covariance matrix R_e .

We will consider a possible open or closed loop scenario in which the input and output data sequences are generated. Also the case of output only identification is considered, making the approach outlined in this paper a general and new framework to identify linear dynamical systems.

The problem considered can be formulated as follows:

Given the input-output (i/o) data batches $\{u(k), y(k)\}_{k=1}^N$ that are generated with a system belonging to the class of LTI systems as represented by (1), the problem is using short length i/o data (with N larger but of the order of n) to estimate the system order, denoted by \hat{n} , and to determine approximate system matrices $(\hat{A}_T, \hat{B}_T, \hat{C}_T, \hat{D}, \hat{K}_T)$ that define the observer:

$$\begin{cases} \hat{x}_T(k+1) = \hat{A}_T \hat{x}_T(k) + \hat{B}_T u_v(k) + \hat{K}_T (y_v(k) - \hat{C}_T \hat{x}_T(k)) \\ \hat{y}_v(k) = \hat{C}_T \hat{x}_T(k) + \hat{D} u_v(k) \end{cases} \quad (2)$$

with $\hat{x}_T(k) \in \mathbb{R}^{\hat{n}}$, such that the approximated output $\hat{y}_v(k)$ is close to the measured output $y_v(k)$ of the validation pair $\{u_v(k), y_v(k)\}_{k=1}^{N_v}$ as expressed by a small value of the cost function,

$$\frac{1}{N_v} \sum_{k=1}^{N_v} \|y_v(k) - \hat{y}_v(k)\|_2^2 \quad (3)$$

The quantitative notions like “small” and “approximate” will be made more precise in the new N2SID solution toward this problem. It should be remarked that contrary to many earlier Subspace Identification (SID) methods, the present problem formulation explicitly takes a prediction error cost function like (3) into consideration.

A key starting point in the formulation of subspace methods is the relation between (structured) Hankel matrices constructed from the i/o data. This relationship will as defined in Verhaegen and Verdult [2007] be the Data Equation. It will be presented in the next section. Here we will also highlight briefly how existing subspace methods have missed to take important structural information about the matrices in this equation into account from the first steps of the solution. N2SID will overcome these shortcomings. Taking these structural properties into consideration makes N2SID attractive especially for identifying MIMO LTI models when only having short data batches.

3. THE DATA EQUATION AND ITS STRUCTURE

Let the LTI model (1) be represented in its so-called observer form:

$$\begin{cases} x(k+1) = (A - KC)x(k) + (B - KD)u(k) + Ky(k) \\ y(k) = Cx(k) + Du(k) + e(k) \end{cases} \quad (4)$$

We will denote this model compactly as:

$$\begin{cases} x(k+1) = \mathcal{A}x(k) + \mathcal{B}u(k) + Ky(k) \\ y(k) = Cx(k) + Du(k) + e(k) \end{cases} \quad (5)$$

with \mathcal{A} the observer system matrix ($A - KC$) and \mathcal{B} equal to $(B - KD)$. Though this property will not be used in the sequel, the matrix \mathcal{A} can be assumed to be asymptotically stable.

For the construction of the data equation, we store the measured i/o data in (block-) Hankel matrices. For fixed N assumed to be larger than the order n of the underlying system, the definition of the number of block-rows fully defines the size of these Hankel matrices. Let this dimensioning parameter be denoted by s , and let $s > n$, then the Hankel matrix of the input (similarly for the output) is defined as:

$$U_s = \begin{bmatrix} u(1) & u(2) & \cdots & u(N-s+1) \\ u(2) & u(3) & & \vdots \\ \vdots & & \ddots & \\ u(s) & u(s+1) & \cdots & u(N) \end{bmatrix} \quad (6)$$

The Hankel matrix defined from the output $y(k)$ and the innovation $e(k)$ are denoted by Y_s and E_s , respectively. The relationship between these Hankel matrices, that readily follows from the linear model equations in (5), require the definition of the following *structured* matrices. First we define the (extended) observability matrix \mathcal{O}_s .

$$\mathcal{O}_s^T = \begin{bmatrix} C^T & \mathcal{A}^T C^T & \cdots & \mathcal{A}^{T^{s-1}} C^T \end{bmatrix} \quad (7)$$

Second, we define a Toeplitz matrix from the quadruple of systems matrices $\{\mathcal{A}, \mathcal{B}, C, D\}$ as:

$$T_{u,s} = \begin{bmatrix} D & 0 & \cdots & 0 \\ C\mathcal{B} & D & & 0 \\ \vdots & & \ddots & \\ C\mathcal{A}^{s-2}\mathcal{B} & \cdots & D \end{bmatrix} \quad (8)$$

and in the same we define a Toeplitz matrix $T_{y,s}$ from the quadruple $\{\mathcal{A}, K, C, 0\}$. Finally, let the state sequence be stored as:

$$X = [x(1) \ x(2) \ \cdots \ x(N-s+1)] \quad (9)$$

Then the data equation compactly reads:

$$Y_s = \mathcal{O}_s X + T_{u,s} U_s + T_{y,s} Y_s + E_s \quad (10)$$

This equation is a simple linear matrix equation that highlights the challenges in subspace identification, which is to approximate from the given Hankel matrices Y_s and U_s the column space of the observability matrix and/or that of the state sequence.

The equation is highly structured. For the sake of brevity in this paper we focus on the following two key structural properties:

- (1) The matrix product $\mathcal{O}_s X$ is *low rank* since $s > n$.
- (2) The matrices $T_{u,s}$ and $T_{y,s}$ are (block-) Toeplitz.

In all existing subspace identification methods, these two key structural matrix properties are *not used* in the first step of the algorithmic solution. Some pre-processing step of the data (Hankel) matrices is usually performed, followed by a low rank factorization. To further illustrate this point, consider the approach in Chiuso [2007] as an example method. Then the first step consists in the estimation of a high order ARX model that is defined by the last (block-) row of the data equation (10) *neglecting*

the term $C\mathcal{A}^{s-1}X$. This is based on the fact that \mathcal{A} is asymptotically stable and the assumption that s is chosen large enough. Then in a later step the estimated ARX parameters are used to define a matrix that asymptotically (both in terms of s and N) is of low rank n . Other recent methods that make use of the application of the nuclear norm, try to enforce the low rank property to an already pre-processed matrix. For example in Liu et al. [2013] the case of measurement noise was considered only, and in a pre-processing step the input Hankel matrix U_s is annihilated from the data equation by an orthogonal projection.

Though statistical consistency generally holds for the existing subspace identification schemes, they refrain from exploiting key structural matrix properties in the data equation in the first step of the algorithm. Such structural information may be key when dealing with short data batches. Therefore in the next section we will revise subspace identification and formulate the new N2SID approach.

4. N2SID

4.1 Pareto optimal Subspace Identification

From the data equation (10) it follows that the minimum variance prediction of the output equals $\hat{y}(k) = y(k) - e(k)$. Let the Hankel matrix \hat{Y}_s be defined from this sequence $\hat{y}(k)$ as we defined Y_s from $y(k)$. Then the data equation becomes:

$$\hat{Y}_s = \mathcal{O}_s X + T_{u,s} U_s + T_{y,s} Y_s \quad (11)$$

Let $\mathcal{T}_{p,m}$ denote the class of lower triangular (block-) Toeplitz matrices with block entries $p \times m$ matrices and let \mathcal{H}_p denote the class of (block-) Hankel matrices with block entries of p column vectors. Then the two key structural properties listed in Section 3 are taken into account in the following problem formulation:

$$\min_{\hat{Y}_s \in \mathcal{H}_p, T_{u,s} \in \mathcal{T}_{p,m}, T_{y,s} \in \mathcal{T}_{p,p}} \text{rank}(\hat{Y}_s - T_{u,s} U_s - T_{y,s} Y_s) \quad (12)$$

and $\min \mathbb{E}[(y(k) - \hat{y}(k))(y(k) - \hat{y}(k))^T]$, where \mathbb{E} denotes the expectation operator. This optimization problem seeks for the Pareto optimal solution with respect to the two cost functions $\text{rank}(\hat{Y}_s - T_{u,s} U_s - T_{y,s} Y_s)$ and $\mathbb{E}[(y(k) - \hat{y}(k))(y(k) - \hat{y}(k))^T]$. This optimization is however not tractable. For that purpose we will develop in the next subsection a *convex relaxation*. This will make it possible to obtain all Pareto optimal solutions using scalarization.

4.2 A convex relaxation

A convex relaxation of the NP hard problem formulation in (12) will now be developed. The original problem is reformulated in two ways. First the $\text{rank}(\cdot)$ operator is substituted by the nuclear norm. The nuclear norm of a matrix X denoted by $\|X\|_*$ is defined as the sum of the singular values of the matrix X . It is also known as the trace norm, the Ky Fan norm or the Schatten norm Liu and Vandenberghe [2010]. This is known to be a good approximation, Fazel et al. [2001], Fazel [2002].

Second the minimum variance criterion is substituted by the following sample average of the trace of the covariance matrix $\mathbb{E}[(y(k) - \hat{y}(k))(y(k) - \hat{y}(k))^T]$:

$$\frac{1}{N} \sum_{k=1}^N \|y(k) - \hat{y}(k)\|_2^2$$

By introducing a scalarization, or regularization, parameter $\lambda \in [0, \infty)$ all Pareto optimal solutions of the convex reformulation of the N2SID problem can be formulated in **one line**.

$$\begin{aligned} & \min_{\hat{Y}_s \in \mathcal{H}_p, T_{u,s} \in \mathcal{T}_{p,m}, T_{y,s} \in \mathcal{T}_{p,p}} \|\hat{Y} - T_{u,s}U_s - T_{y,s}Y_s\|_* + \\ & \frac{\lambda}{N} \sum_{k=1}^N \|y(k) - \hat{y}(k)\|_2^2 \end{aligned} \quad (13)$$

It is well-known that this problem can be recast as a semi-definite programming problem, Fazel et al. [2001], Fazel [2002], and hence it can be efficiently solved with standard solvers. We have used the modeling language YALMIP, Löfberg [2001], to perform experiments, results of which we will present next.

The method encompasses in a straightforward manner the identification problems with output data only. In that case the convex relaxed problem formulation reads:

$$\begin{aligned} & \min_{\hat{Y}_s \in \mathcal{H}_p, T_{y,s} \in \mathcal{T}_{p,p}} \|\hat{Y} - T_{y,s}Y_s\|_* + \\ & \frac{\lambda}{N} \sum_{k=1}^N \|y(k) - \hat{y}(k)\|_2^2 \end{aligned} \quad (14)$$

5. ILLUSTRATIVE EXAMPLES

5.1 Open-loop experiment

In this section we report results on numerical experiments in Matlab. In the first example we consider 100 second order single-input single-output systems as in (1) randomly generated with the command `drss`. The matrix K has been generated with the command `randn` which gives a matrix of elements drawn from a standardized normal density function. Any model for which the absolute value of the largest eigenvalue of the system matrix A is larger than 0.99 has been discarded. Data for system identification has been generated for each model and with time horizon $N = 50$. The noise $e(k)$ is white and drawn from a normal density function with standard deviation equal to 0.2. The input signal $u(k)$ is a sequence of ± 1 obtained by taking the sign of a vector of values obtained from a standardized normal density function. The initial value is obtained from a normal density function with standard deviation 5, where the components are uncorrelated. The parameter s has been equal to 15. We do not consider the nuclear norm of the matrix as defined in (13), but instead we first multiply it with a random matrix from the right. This random matrix was obtained from a standardized normal density function and the number of columns was 22. It is known that this type of randomization is a powerful tool in low rank matrix approximation, Halko et al. [2011]. The reason we do this modification is that it will reduce the computational complexity without significantly affecting the quality of the results obtained. We choose model order by

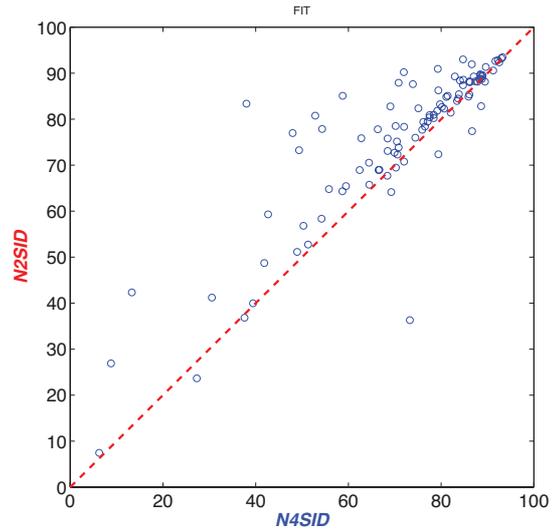


Fig. 1. Scatter plot showing the fit N2SID versus N4SID for 100 randomly generated examples.

looking at the singular values of the matrix that minimizes the nuclear norm. The model order is equal to the index of the singular values that is closest to the logarithmic mean of the largest and smallest singular value. However, in case that index is greater than 10 we choose as model order 10.

Since our approach gives as solution a whole family of models parameterized with regularization parameter λ we perform a second optimization over this parameter with respect to a fit criterion as implemented in the command `FIT` of the System Identification Toolbox in Matlab. We consider 20 logarithmically spaced values of λ/N in the interval $[10^{-0.5}, 10^4]$.

We then compare our method with a standard subspace method as implemented in the command `n4sid` of the System Identification Toolbox of Matlab. This algorithm will pick model order equal to the one in the interval 0 to 10 which give the best fit on the identification data. We also make sure that `n4sid` uses the same value of s as we do. We also insist that the direct term in the model is estimated. The comparison between our model and the one of `n4sid` is done by computing the fit of the two models on a second validation data set. In Figure 1 the fit of N2SID versus N4SID is plotted for the 100 random examples. In this plot one negative fit value has been removed. In more than 80% of the cases N2SID has a higher fit value than N4SID. The average fit values are 76.0 for N2SID and 71.1 for N4SID. The average model orders are 6.67 for N2SID and 5.01 for N4SID. This clearly demonstrates the advantage of using N2SID on short data sets. We have also performed experiments on longer data sets. Preliminary results show that there is no significant difference between the two methods. This is an indication that for large data length batches both methods deliver consistent results.

5.2 Closed-loop experiment

In the second example we consider closed loop identification. For this purpose we consider a model on the form (1), where

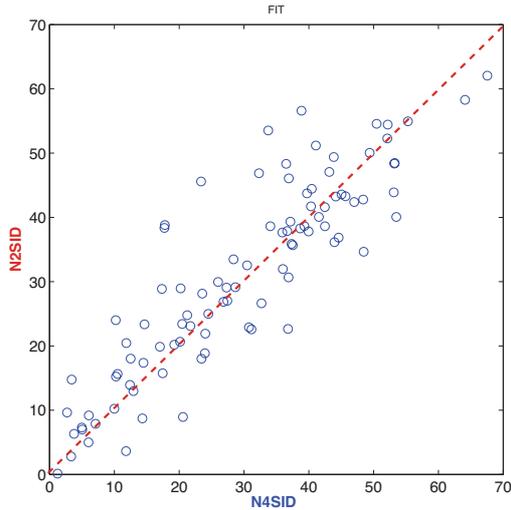


Fig. 2. Scatter plot showing the fit N2SID versus N4SID for 100 different realizations of the reference value and the noise.

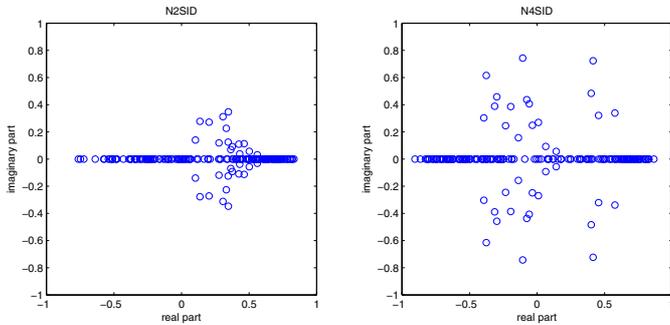


Fig. 3. Scatter plot showing the eigenvalues of the A -matrix for N2SID and N4SID for 100 different realizations of the reference value and the noise.

$$A = \begin{bmatrix} 0 & 1 \\ 0 & 0.7 \end{bmatrix}; \quad B = \begin{bmatrix} 0 \\ 1 \end{bmatrix}; \quad K = \begin{bmatrix} -0.3 \\ 0.04 \end{bmatrix}$$

$$C = [1 \ 0]; \quad D = 0$$

We control the model with an observer-based state feedback where K is the observer gain and where the state feedback matrix is $L = [0.25 \ -0.3]$. This design will place the closed loop eigenvalues all at 0.5. The controller also has a feed-forward from a reference value $r(k)$, which is a sequence of ± 1 obtained by taking the sign of a vector of values obtained from a standardized normal density function. This signal is multiplied with a gain before it is added to the input signal such that the closed loop steady gain from reference value to $y(k)$ is equal to one. The noise $e(k)$ is white and drawn from a normal density function with standard deviation equal to 0.1. In this experiment we will not look for the best model order, but instead we just compute models of order $n = 2$. All the other parameters are the same as in the previous experiment. We show in Figure 2 the fit of N2SID versus N4SID for 100 different realizations of the reference value and the noise. The fit is better for N2SID in 59 % of the cases. Moreover, as is seen in Figure 3, the spread of the eigenvalues of the A -matrix is smaller for N2SID as compared to N4SID.

6. CONCLUDING REMARKS

Subspace identification is revisited in this paper in the scope of nuclear norm optimization methods. A new way to impose structural matrix properties in the data equation in subspace identification on the measured data has been presented. The new subspace identification method is referred to as N2SID. It is shown that especially for small data length batches when the number of samples is only a small multiple of the order of the underlying system, the incorporation of structural information about the low rank property of the matrix revealing the required subspace and the (block) Toeplitz structure of the matrix containing the unknown Markov parameters enables to improve the results of widely used SID methods. In addition to the structural constraints the N2SID method also enables to make a trade-off in the first step of the calculations between the subspace approximation and the prediction error cost function. As such it also overcomes the persistent drawback that SID did not consider a (classical prediction error) cost function.

The single integrative step that aims at imposing key structural matrix properties makes a trade-off between the prediction error cost function and the problem to retrieve the subspace of interest. This integrative approach may help to simplify the analysis of the optimality of SID methods and to further clarify their link with prediction error methods. This new way of looking upon SID will open up the possibility for new developments in the future.

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